

Amendments to the Claim:

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

1-11 (cancelled).

12 (currently amended). An isolated polypeptide which ~~comprising an amino acid sequence having at least 75% sequence identity with a reference sequence which is SEQ ID NO:2, or a fragment thereof,~~

- (a) comprises SEQ ID NO:2;
- (b) consists of residues 234-1791 of SEQ ID NO:2;
- (c) differs from the polypeptide of (b) solely by deletion of 1-10 residues from the amino terminal of (b) and/or deletion of 1-10 residues from the carboxy terminal of (b);
- (d) differs from (b) solely by addition of 1-10 residues to the amino terminal or carboxy terminal of (b); or
- (e) is at least 97% identical to a polypeptide of (a), (b), (c), or (d)

~~wherein said polypeptide, and, if said reference sequence is a fragment, said fragment,~~

- i) has a proteolytic activity specific at least for Insulin Like Growth Factor Binding Protein 5 (IGFBP-5); and/or
- ii) is recognised by an antibody, or a binding fragment thereof, which recognizes a polypeptide having the amino acid sequence as shown in SEQ ID NO:2; and/or
- iii) competes with a polypeptide having the amino acid sequence as shown in SEQ ID NO:2 for binding to a cell surface receptor with an affinity for said

polypeptide.

13-16 (cancelled).

17 (currently amended). The polypeptide of Polypeptide ~~according to~~ claim ~~14~~ 12, wherein the prepro part of PAPP-A2 is operably linked to the mature part of PAPP-A2 corresponding to amino acid residues 234 to 1791 of SEQ ID NO:2.

18. (currently amended). The polypeptide of Polypeptide ~~according to~~ claim 12, wherein said polypeptide is a recombinant polypeptide.

19. (currently amended). The polypeptide of Polypeptide ~~according to~~ claim 12, wherein the polypeptide is free of human proteins, or other proteins ~~natively~~ naturally associated with said polypeptide.

20-29 (cancelled).

30 (previously presented). A method for detecting PAPP-A2, or measuring the level of PAPP-A2, in a biological sample obtained from an individual, said method comprising the steps of

- i) obtaining a biological sample from said individual,
- ii) detecting PAPP-A2 in said sample by detecting a polypeptide according to claim 12.

31 (original). Method of claim 30, said method comprising the further step of comparing the PAPP-A2 or the level of PAPP-A2 detected in step ii) with a predetermined value selected from the group consisting of

- i) a predetermined amount and/or concentration of PAPP-A2; and/or
- ii) a predetermined amount and/or concentration of PAPP-A2 mRNA; and/or
- iii) a predetermined PAPP-A2 specific protease activity.

32 (original). Method of claim 31, wherein said predetermined value is indicative of a normal physiological condition of said individual.

33 (original). The method of claim 30, wherein said biological sample is selected from the group consisting of blood, urine, pleural fluid, oral washings, tissue biopsies, and follicular fluid.

34 (original). The method of claim 30, wherein said level of PAPP-A2 is measured as PAPP-A2 specific protease activity.

35 (original). The method of claim 30, wherein said level of PAPP-A2 is measured as amount of PAPP-A2 protein.

36 (original). The method of claim 30, wherein said level of PAPP-A2 is measured as amount of PAPP-A2 messenger RNA.

37 (original). The method of claim 35, wherein said amount of PAPP-A2 protein is measured by immunochemical analysis.

38 (original). The method of claim 37, wherein said amount of PAPP-A2 protein is detected by at least one monoclonal antibody.

39 (original). The method of claim 30, wherein said PAPP-A2 protein is detected in a complex comprising at least one additional component, preferably a polypeptide.

40 (original). The method of claim 30, wherein said PAPP-A2 is detected as a PAPP-A2 monomer.

41 (original). The method of claim 30, wherein said PAPP-A2 is detected as a PAPP-A2 dimer.

42 (previously presented). A method of diagnosing a clinical condition in an individual, said method comprising the steps of

- i) performing the method of claim 30, and
- ii) diagnosing the clinical condition.

43 (original). Method of claim 42, wherein said clinical condition is a fetal abnormality.

44 (original). The method of claim 43, wherein said fetal abnormality is selected from the group consisting of Trisomy 21, Trisomy 18, Trisomy 13, and Open Spina Bifida.

45 (original). The method according to claim 43, wherein

said fetal abnormality is ectopic pregnancy, open spina bifida, neural tube defects, ventral wall defects, Edwards Syndrome, Patau's Syndrome, Turner Syndrome, Monosomy X or Klinefelter's Syndrome.

46 (original). The method of claim 43, wherein said clinical condition is an altered growth state selected from the group consisting of a growth promoting state and a growth inhibiting state.

47 (original). The method of claim 46, wherein said clinical condition is selected from the group consisting of restenosis, atherosclerosis, wound healing, fibrosis, myocardial infarction, osteoporosis, rheumatoid arthritis, multiple myeloma, or cancer.

48 (cancelled).

49 (previously presented). A method for identifying an agent inhibiting the protease activity of PAPP-A2, said method comprising the steps of

- i) incubating a) the polypeptide according to claim 12 and b) a predetermined substrate for said polypeptide, and c) a putative inhibitory agent, and
- ii) determining if proteolysis of said substrate is inhibited.

50 (original). The method of claim 49, wherein said substrate comprises a polypeptide.

51 (original). The method of claim 50, wherein said substrate comprises an internally quenched fluorescent peptide.

52 (original). The method of claim 50, wherein said substrate comprises or essentially consists of IGFBP-5, or a fragment thereof.

53-54 (cancelled).

55 (currently amended). A method for identifying an agent enhancing the protease activity of PAPP-A2, said method comprising the steps of

- i) incubating a) the polypeptide according to claim 12 and b) a predetermined substrate for said polypeptide,

and c) a putative enhancer agent, and

- ii) ~~determinining~~ determining if proteolysis of said substrate is enhanced.

56 (previously presented). The method of claim 55, wherein said substrate comprises a polypeptide.

57 (previously presented). The method of claim 55, wherein said substrate comprises an internally quenched fluorescent peptide.

58 (previously presented). The method of claim 55, wherein said substrate comprises or essentially consists of IGFBP-5, or a fragment thereof.

59-61 (cancelled).

62 (currently amended). A method for purification of PAPP-A2 or complexes of PAPP-A2 with other proteins, said method comprising the steps of

- i) ~~provising~~ providing a polyclonal or monoclonal antibody with specific binding affinity for a polypeptide according to claim 12,
- ii) purifying PAPP-A2 by means of affinity chromatography.

63-69 (cancelled).

70 (previously presented). The polypeptide of claim 12 wherein said polypeptide is the polypeptide according to (a), 9b), (c) or (d) SEQ ID NO:2, or a fragment thereof.

71-74 (cancelled).

75 (currently amended). The polypeptide of claim 12 wherein said polypeptide is the polypeptide according to (e) comprising an amino acid sequence having at least 98% sequence identity with said reference sequence.

76-82 (cancelled).

83 (currently amended). The polypeptide of claim 12 which wherein said polypeptide comprises an amino acid sequence which differs from said ~~reference~~ sequence (b) solely by one or more conservative substitutions.

84 (cancelled).

85 (currently amended). The polypeptide of claim 12 wherein said polypeptide ~~which~~ comprises an amino acid sequence which is identical to amino acids 234 to 1791 of SEQ ID NO:2.

86 (cancelled).

87 (currently amended). The polypeptide of claim 12 wherein said polypeptide ~~which~~ consists of amino acids 234 to 1791 of SEQ ID NO:2.

88-89 (cancelled).

90 (new). An isolated polypeptide which is

- (1) a polypeptide consisting of an amino acid sequence which is (a) identical to residues 234-1791 of SEQ ID NO:2, or (b) a fragment, at least 6 amino acids in length, of mature PAPP-A2 (residues 234-1791 of SEQ ID NO:2)

where said fragment

- i) has a proteolytic activity specific at least for Insulin Like Growth Factor Binding Protein 5 (IGFBP-5); and/or
- ii) is recognized by an antibody, or a binding fragment thereof, which recognizes a polypeptide having the amino acid sequence as shown in SEQ ID NO:2; and/or
- iii) competes with a polypeptide having the amino acid sequence as shown in SEQ ID NO:2 for binding to a cell surface receptor with an affinity for said polypeptide

or

- (2) a polypeptide which consists of a fusion of the polypeptide of (1) with an immunogenic carrier protein, or with a tag which may be used to facilitate the detection or purification of the fusion, with the proviso that said fusion of (2) is not a pregnancy-associated plasma protein.

91 (new). The polypeptide of claim 90 which is a fragment according to (1).

92 (new). The polypeptide of claim 91 which is a fragment, at least 17 amino acids in length, of mature PAPP-A2.

93 (new). The polypeptide of claim 91 which comprises at least 75% of mature PAPP-A2.

94. The polypeptide of claim 91 which is a fragment comprising at least one of the following regions of SEQ ID NO:2:

Cys-403 to Cys-499

Cys-828 to Cys-881

Cys-1048 to Cys-1115

Cys-1390 to Cys-1396

Cys-1459 to Cys-1464

Cys-1521 to Cys-1525

Cys-1590 to Cys-1595

Cys-1646 to Cys-1653

Cys-1729 to Cys-1733.

95 (new). The polypeptide of claim 91 which comprises the elongated zinc binding consensus sequence, LNR 1-3, SCR 1-5, and all cysteine residues of mature PAPP-A2.

96 (new). The polypeptide of claim 90 where the fragment is at least 50 AA in length.

97 (new). An isolated polypeptide comprising amino acids 1-22 of SEQ ID NO:2.

98 (new). An isolated polypeptide comprising amino acids 23-233 of SEQ ID NO:2.

99 (new). An isolated polypeptide comprising amino acids 1-233 of SEQ ID NO:2.

100 (new). The polypeptide of claim 91 which is a processing variant of mature PAPP-A2.